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(FILE 'HOME' ENTERED AT 07:39:58 ON 12 JAN 2007)

INDEX 'ADISCTI, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, ANTE, AQUALINE, AQUASCI, BIOENG, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CAPLUS, CEABA-VTB, CIN, CONFSCI, CROPB, CROPU, DDFB, DDFU, DGENE, DISSABS, DRUGB, DRUGMONOG2, DRUGU, EMBAL, EMBASE, ...' ENTERED AT 07:40:13 ON 12 JAN 2007
SEA AMYLASE

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366 FILE ADISNEWS
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 2484 FILE USPAT2
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 247 FILE VETU
 72 FILE WATER
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 60 FILE WPIFV
 8563 FILE WPINDEX

L1 QUE AMYLASE

FILE 'CAPLUS, BIOSIS, MEDLINE, SCISEARCH, EMBASE, PASCAL, CABA,
 TOXCENTER, WPIDS, FSTA, JICST-EPLUS, BIOTECHDS, AGRICOLA, ESBIODBASE,
 LIFESCI, BIOTECHNO! ENTERED AT 07:41:25 ON 12 JAN 2007

L2 9502 S L1 AND (MUTANT OR VARIANT)
 L3 0 S L2 AND (ASP128, GLY140, SER144,)
 L4 4 S L2 AND (POSITION 128)
 L5 1 DUP REM L4 (3 DUPLICATES REMOVED)
 L6 0 S L2 AND (POSITION 140, POSITION 144, POSITION 168, POSITION 181
 L7 0 S L2 AND (AMINO ACID 140)
 L8 0 S L2 AND (ASPL28, GLYL40, SER144, ARG168, ASNL8L, GLU207, PHE2

=> d 15 ibib ab

L5 ANSWER 1 OF 1 CAPLUS COPYRIGHT 2007 ACS on STN DUPLICATE 1
ACCESSION NUMBER: 2000:842252 CAPLUS
DOCUMENT NUMBER: 134:14747
TITLE: Subtilase enzymes of the I-S1 and I-S2 subgroups
having at least one additional amino acid residue
between **positions 128** and 129 with
improved wash performance
INVENTOR(S): Vilbour, Andersen Kim; Mikkelsen, Frank F.; Kamp,
Hansen Peter; Norregaard-Madsen, Mads
PATENT ASSIGNEE(S): Novo Nordisk A/S, Den.
SOURCE: PCT Int. Appl., 68 pp.
CODEN: PIXXD2
DOCUMENT TYPE: Patent
LANGUAGE: English
FAMILY ACC. NUM. COUNT: 20
PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 2000071690	A1	20001130	WO 2000-DK241	20000510
W:	AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, DZ, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW			
RW:	GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG			
EP 1183342	A1	20020306	EP 2000-925092	20000510
R:	AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, SI, LT, LV, FI, RO			

PRIORITY APPLN. INFO.: DK 1999-712 A 19990520
WO 2000-DK241 W 20000510

AB Subtilase enzymes of the I-S1 and I-S2 sub-groups are provided having an addnl. amino acid residue in **position 128** of the active site loop (c) region from positions 125 to 132. Specifically, Savinase (Bacillus lentus subtilisin 309) **variants** are provided having at least one amino acid residue inserted in between **positions 128** and 129 (numbering based on subtilisin BPN'). Site-specific mutagenic primers are used to insert the desired codons at the desired position(s) of the wild-type gene, and the **mutant** genes are used to transform Escherichia coli or Bacillus subtilis for the fermentation and purification of **variant** subtilisins. These **variant** subtilases exhibit improved wash performance in a detergent in comparison to its parent enzyme. Although this finding was done in subtilisin 309, it is predicted that it will be possible to produce or isolate similar advantageous subtilases or subtilase **variants**. The invention further relates to genes coding for the expression of said enzymes when inserted into a suitable host cell or organism, host cells transformed therewith and capable of expressing said enzyme **variants**, and methods for producing the novel enzymes.

REFERENCE COUNT: 7 THERE ARE 7 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

=> s 12 and (position 140, position 144, position 168, position 181, position 207, position 272, position 375, position 434, position 466)

12 FILES SEARCHED...

L6 0 L2 AND (POSITION 140, POSITION 144, POSITION 168, POSITION 181, POSITION 207, POSITION 272, POSITION 375, POSITION 434, POSITION 466)

=> s l2 and (amino acid 140)

1 FILES SEARCHED...

12 FILES SEARCHED...

L7 0 L2 AND (AMINO ACID 140)

=> s l2 and (Aspl28, Glyl40, Ser144, Arg168, Asnl8l, Glu207, Phe272, Ser375,
Trp434, Glu466)

L8 0 L2 AND (ASPL28, GLYL40, SER144, ARG168, ASNL8L, GLU207, PHE272,
SER375, TRP434, GLU466)

Refine Search

Search Results -

Term	Documents
ASPL28	0
ASPL28S	0
GLYL40	0
GLYL40S	0
SER144	3
SER144S	0
ARG168	11
ARG168S	0
ASNL8L	0
ASNL8LS	0
GLU207	4
(ASPL28, GLYL40, SER144, ARG168, ASNL8L, GLU207, PHE272, SER375, TRP434, GLU466).PGPB,USPT,USOC,EPAB,JPAB,DWPI.	0

There are more results than shown above. [Click here to view the entire set.](#)

Database:

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US OCR Full-Text Database
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Derwent World Patents Index
IBM Technical Disclosure Bulletins

Search:

L2 and (position 128, position 140,
position 144, position 168, position
181, position 207, position 272,



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Query

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Count

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Name

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result set

DB=PGPB,USPT,USOC,EPAB,JPAB,DWPI; PLUR=YES; OP=ADJ

<u>L5</u>	Aspl28, Glyl40, Ser144, Arg168, Asnl8l, Glu207, Phe272, Ser375, Trp434, Glu466	0	<u>L5</u>
<u>L4</u>	L3 and (Aspl28, Glyl40, Ser144, Arg168, Asnl8l, Glu207, Phe272, Ser375, Trp434 Glu466)	0	<u>L4</u>
<u>L3</u>	L2 same (variant or mutant)	4873	<u>L3</u>
<u>L2</u>	amylase	32511	<u>L2</u>
<u>L1</u>	amylase	32511	<u>L1</u>

END OF SEARCH HISTORY

Refine Search

Search Results -

Term	Documents
AMYLASE	27945
AMYLASES	8961
SER375	3
SER375S	0
(SER375 SAME AMYLASE).PGPB,USPT,USOC,EPAB,JPAB,DWPI.	0
(AMYLASE SAME SER375).PGPB,USPT,USOC,EPAB,JPAB,DWPI.	0

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L20

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Name
 result set

DB=PGPB,USPT,USOC,EPAB,JPAB,DWPI; PLUR=YES; OP=ADJ

L20 amylase same Ser375

0 L20

L19 amylase same Trp434

0 L19

L18 amylase same Glu466

0 L18

L17 (amylase same Glu466).clm.

0 L17

L16 (amylase same Trp434).clm.

0 L16

<u>L15</u>	(amylase same Ser375).clm.	0	<u>L15</u>
<u>L14</u>	(amylase same Phe272).clm.	0	<u>L14</u>
<u>L13</u>	(amylase same Glu207).clm.	1	<u>L13</u>
<u>L12</u>	(amylase same Asn 181).clm.	0	<u>L12</u>
<u>L11</u>	(amylase and Arg 168).clm.	0	<u>L11</u>
<u>L10</u>	(amylase and Ser 144).clm.	0	<u>L10</u>
<u>L9</u>	(amylase and Gly140).clm.	0	<u>L9</u>
<u>L8</u>	(amylase and Asp 128).clm.	0	<u>L8</u>
<u>L7</u>	L2 and (position 128)	251	<u>L7</u>
<u>L6</u>	L2 and (position 144)	341	<u>L6</u>
<u>L5</u>	Asp128, Gly140, Ser144, Arg168, Asn181, Glu207, Phe272, Ser375, Trp434, Glu466	0	<u>L5</u>
<u>L4</u>	L3 and (Asp128, Gly140, Ser144, Arg168, Asn181, Glu207, Phe272, Ser375, Trp434 Glu466)	0	<u>L4</u>
<u>L3</u>	L2 same (variant or mutant)	4873	<u>L3</u>
<u>L2</u>	amylase	32511	<u>L2</u>
<u>L1</u>	amylase	32511	<u>L1</u>

END OF SEARCH HISTORY